3. L'Rumant



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Page 1 of 7

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## TECH CENTER 1600/2900

RAW SEQUENCE LISTING | DATE: 11/07/2000
PATENT APPENDATION: US/09/230,111B | TIME: 1::01:2

Imput Set: A:\48962ap1.app Ont.put Set: N:\CRF3\11072000\1230111B.raw

```
3 HITUS APPLICANT: Sato, Taka-Aki
             Yanagisawa, June
                                                                                     Does Not Comply
      \delta \sim 1200 fills of invention: Compounds that inhibit interaction between
                                                                                Corrected Diskette Needed
             SIGNAL-TRANSDUCING PROTEIRS AND THE GLOF (FOL/DHR)
              DOMAIN AND USES THEFEOF
     10 ×130× FILE REFERENCE: 48962-A-POT-US
     12 STAUS CORRENT APPLICATION NUMBER: 09/230,411B
     13 -141> CURRENT FILING DATE: 1999-05-17
     15 KINDS HUMBER OF SPO 1D NOS: 33
     17 <170> SOFTWARE: PatentIn Ver. 2.1
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     30 <222> LOCATION: (1)
     31 <223> OTHER INFORMATION: Xaa=Gly, Ser, Ala or Glu
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     35 <222> LOCATION: (4)
     36 <223> OTHER INFORMATION: Xaa-Phe. 11e or Leu
    38 4400> SEQUENCE; 1
W--> 39 Xaa Leu Gly Xaa
    40 1
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    55~<\!\!223\!\!> OTHER INFORMATION: Xaa-Lys, Arg or Gln
    57 <220> FEATURE:
                                                           Xaa may only represent one residue. See #6
    58 <221> NAME/KEY:
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    59 <222> LOCATION: (2)
    60 <223> OTHER INFORMATION: (Xaa-any 2-4 amino acids
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    63 <221> NAME/KEY: SITE
    64 <222> LOCATION: (3)
                                                            on the Error Summery Sheet
    65\ <\!\!223\!\!> OTHER INFORMATION: Maa-Gly, Sec. Ala or Glu
```



Imput Ser : A:\48962apl.app

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```
67 KE200 FEATURE:
     68 .231> NAME/KEY, SITE
     60 222× 100AT16N: (6)
     20 -2250 OTHER INFORMATION: XeasPho, lie or Lou
72 1400 - SEMBERGE: 2
W--> 73 Xaa Xaa Xaa Leu Gly Xaa
                                    See p./
     77 H2105 SEQ 10 NO: 3
     78 -2115 LENGTH: 4
     79 -. 212> PYPE: PRT
     80 -2:3> ORGANISM: Artificial Sequence
     83 H220> FEATURE:
     83 -223> OTHER INFORMATION: Description of Artificial
            Sequence:source:synthesized
     86 +460> SEQUENCE: 3
     87 Ser Leu Gly Tle
     88 1
     91 -.210> SEQ ID NO: 4
     92 - 2115 LENGTH: 3
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     94 <\!\!(21.3\!\!> ORGANISM: Arrificial Sequence
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     108 <223> OTHER INFORMATION: Xaa=any one amino acid
     110 <220> FEATURE:
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     112 <222> LOCATION: (3)
    113 <223> OTHER INFORMATION: Xaa*Val, He or Leu
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    117 1
    120 <210> SEQ ID NO: 5
    121 <211> GENGTH: 15
    122 <212> TYPE: PRT
   1123 4213> ORGANISM: human
    125 <400> SEQUENCE: 5
    126 Asp Ser Glu Ash Ser Ash Phe Arg Ash Glu Lie Gln Ser Leu Val
    127 1
                          5
                                            3.0
    130 <210> SEO ID NO: 6
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    132 <212> TYPE: PRT
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RAW SEQUENCE LISTING DATE: 11/67/2006
PATENT / PPLICETION: US/09/230,111B TIME: 15:01:26

Inpn/ Sct : A:\48962ap1.app
cm:pnc Set: N:\CRF3\11072000\1230111B.raw

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133 K2138 OPGARISM: Call
135 - 400× Sequence: 5
The Ser lie Ser ash Ser Arg Ash Sin Ash Gin Gly Gin Ser Lea Gin
137 1
146 82102 SEC 16 BOL 7
141 -0115 LENGIE: 45
140 C2122 TYPE: PRI
143 <213> ORGANISM: mouse
115 <400> SEQUENCE: 7
145 Sor the Pro Asp The Giy was Gie Ash Gie Gly Cin Cys Lee Gie
                             1.0
150 <210> SPO 10 RE: 8
151 <2.11> LENGIR: 1
152 (212> TYPE: PPT
15% <21% ORGANISM: Artificial Sequence
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157
     Sequence:source:synthesized
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160 Glu Ser Leu Val
161 1
164 <210> SEQ ID NO: 9
165 <211> LFNGTH: 6
166 <212> TYPE: PRT
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Description of Artificial Sequence:
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173 <406> SEQUENCE: 9
174 Thr Tle Gla Ser Val 11e
175 1
178 <210> SEQ 10 NO: 10
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188 Arg Gly Phe Tie Ser Ser Leu Val
189 1
                    5
192 <210> SEO ID NO: 11
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RAW SEQUENCE LISTING DATE: 11/07/2000 PATENT APPLICATION: US/09/230,111B TIME: 15:91:29

Imput Set : A:\48962apl.app

output Set: N:\CRF3\11072000\1230111B.raw

```
20% Arg Glo Tur lie din Ser The Val
203 1
200 (210) SEC 35 NG: 12
207 (21) EMNOTE: 11
208 (212) TYPE: PRU
200 - 213> ORGANISH: Artificial Sequence
DIT CHEUN PEATURE:
212 2232 OTHER INFORMATION: Description of Artificial
        Sequence:source:synthesized
213
215 -400> SEQUENCE: 12
216 din Ash Phe Ary Th: Tyr fle Val Ser Phy Val
2:0
200 4230> SEO 10 NO: 13
221 211> LENGTH: 13
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323 <213> ORGANISM: Artificial Sequence
225 | 220> FEATURE:
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230 Ser Asp Ser Ash Met Ash Met Ash Glu Leu Ser Glu Val
23.1 1.
                     5
                            10
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235 <211> LENGTH: 15
236 <212> TYPE: PRT
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239 - 220> FEATURE:
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2.11
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244 Pro Pro Thr Cys Ser Gln Ala Asa Ser Gly Arg Ile Ser Thr Leu 245 1 5 . 10 15
248 <210> SEQ ID NO: 15
249 <211> LENGTH: 15
250 ×212> TYPE: PRT
251 <213> ORGANISM: Artificial Sequence
253 -220> FEATURE:
254 <223> OTHER INFORMATION: Description of Artificial
255
        Sequence:source:synthesized
257 <400> SEQUENCE: 15
258 The Asp Leu Ala Ser Glu Phe Leu Phe Leu Ser Asm Ser Phe Leu
259 1
             5
                          10
262 <210> SEQ ID NO: 16
263 <211> LENGTH: 15
264 <212> TYPE: PRT
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267 <220> FEATURE:
268 <223> OTHER INFORMATION: Description of Artificial
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## RAW SEQUENCE LISTING DATE: 41/07/2000 PATENT APPLICATION: US/09/230,111B TIME: 15:01:29

Imput Set : A:\48962apl.app ontput Set: N:\CRF3\11072000\1230111B.raw

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271 <4000 SEJUENCE: 16
272~\mathrm{Asp} Ser Glu Het \mathrm{T}_{24} Ash Phe Arg Ser Glu Lou Ala Ser Val Val
                                          3.3
276 <2105 SEQ 1D NO: 17
277 -271> LENGTH: 15
278 ×212> PYPE: PRI
279 <213> ORGANISM: Aftificial Sequence
281 - 220> FEATURE:
282 /223> OTHER INFORMATION: Description of Artificial
283
         Sequence:source:synthesized
285 .100 - SEQUENCE: 17
286 The Pro Pro Asp Ser Glu Asp Gly Asu Glu Glu Glu Ser Leu Vol
287 .1
                                           10
290 2105 SEQ (D NO: 18
294\ \text{\ensuremath{$\sim$}}\ 2115\ \text{LENGTH:}\ 4
292 <212> TYPE: PRT
293 <213> ORGANISH: Artificial Sequence
295 <220> FEATURE:
296 -223> OTHER INFORMATION: Description of Artificial
297
         Sequence:source:synthesized
299 <400> SEQUENCE: 18
300 Gin Ser Leu Val
201 1
304 <210> SEO ID NO: 19
305 <211> LENGTH: 5
306 <212> TYPE: PRT
307 <213> ORGANISM: Artificial Sequence
309 (220> FEATURE:
310 <2235 OTHER INFORMATION: Description of Artificial Sequence: source
311
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313 <400> SEQUENCE: 19
314 Ile Gin Ser Leu Val
315 1
316 <210> SEQ ID NO: 20
319 <211> LENGTH: 6
320 <212> TYPE: PRT
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
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325
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327 <400> SEQUENCE: 20
328 Glu Ile Gln Ser Leu Val
329 1
332 <210> SEQ ID NO: 21
333 <211> LENGTH: 7
334 <21.2> TYPE: PRT
335 <213> ORGANISM: Artificial Sequence
337 <220> PEATURE:
338 <223> OTHER INFORMATION: Description of Artificial
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VERIFICATION SUMMARY PATERT ARELICATION: US/09/230,111B

DATE: 11/01/2003 [FBB: 15:01:30

Impu: Set : A:\48962apl.app

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1:73 H:341 w: (46) "n" or "Xna" used, for SEO 19#:2
1:116 M:341 w: (46) "n" or "Xna" used, for SEO 10#:3
1:106 M:258 w: Mandatory feature missing, 221% not fermi for SEO 10#:3
1:10% M:258 w: Mandatory reature missing, 222% not found for SEO 10#:3
1:10% M:258 w: Mandatory reature missing, 222% not found for SEO 10#:3
1:10% M:258 W: Mandatory Feature missing, 221% not feund for SEO 10#:3
1:1623 M:258 W: Mandatory Feature missing, 222% not found for SEO 10#:3
1:1623 M:258 W: Mandatory Feature missing, 222% not found for SEO 10#:3
1:1640 M:258 W: Mandatory Feature missing, 221% not found for SEO 10#:3
1:1640 M:258 W: Mandatory Feature missing, 221% not found for SEO 10#:3
1:1640 M:258 W: Mandatory Feature missing, 221% not found for SEO 10#:3
1:1640 M:258 W: Mandatory Feature missing, 221% not found for SEO 10#:33